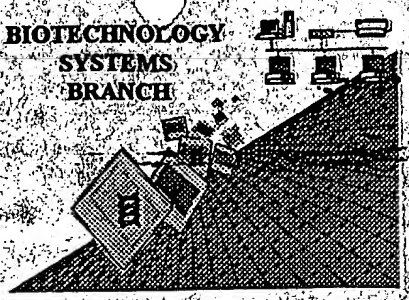
BIOTECHNOLOGY SYSTEMS BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/659,860

Source: OIPE

Date Processed by STIC: 9-22-00

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY**

or,

2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR FURTHER INFORMATION, PLEASE TELEPHONE MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/659,860

DATE: 09/22/2000
TIME: 14:48:08

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

Does Not Comply
Corrected Diskette Needed

```
      3 <110> APPLICANT: Hong Zhang
      4       Andrew T. Watt
      6 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 7 EXPRESSION
      8 <130> FILE REFERENCE: RTS-0201
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/659,860
C--> 10 <141> CURRENT FILING DATE: 2000-09-11
     10 <160> NUMBER OF SEQ ID NOS: 174
     13 <210> SEQ ID NO: 1
     14 <211> LENGTH: 20
     15 <212> TYPE: DNA
     16 <213> ORGANISM: Artificial Sequence
W--> 17 <220> FEATURE:
     18 <223> OTHER INFORMATION: Antisense Oligonucleotide
     20 <400> SEQUENCE: 1
     21 tccgtcatcg ctcctcaggg
                                                                    20
     24 <210> SEQ ID NO: 2
     25 <211> LENGTH: 20
     26 <212> TYPE: DNA
     27 <213> ORGANISM: Artificial Sequence
W--> 29 <220> FEATURE:
     29 <223> OTHER INFORMATION: Antisense Oligonucleotide
     31 <400> SEQUENCE: 2
     32 atgcattctg cccccaagga
     35 <210> SEQ ID NO: 3
     36 <211> LENGTH: 2309
     37 <212> TYPE: DNA
     38 <213> ORGANISM: Homo sapiens
     40 <220> FEATURE:
     41 <221> NAME/KEY: CDS
     42 <222> LOCATION: (44)...(955)
     44 <400> SEQUENCE: 3
     45 gagagactgt gccagtccca gccgccctac cgccgtggga acg atg gca gat gat      55
     46                                                 Met Ala Asp Asp
     47                                                   1
     49 cag ggc tgt att gaa gag cag ggg gtt gag gat tca gca aat gaa gat     103
     50 Gln Gly Cys Ile Glu Glu Gln Gly Val Glu Asp Ser Ala Asn Glu Asp
     51   5                  10                  15                  20
     53 tca gtg gat gct aag cca gac cgg tcc tcg ttt gta ccg tcc ctc ttc     151
     54 Ser Val Asp Ala Lys Pro Asp Arg Ser Ser Phe Val Pro Ser Leu Phe
     55                  25                  30                  35
     57 agt aag aag aag aaa aat gtc acc atg cga tcc atc aag acc acc cgg     199
     58 Ser Lys Lys Lys Lys Asn Val Thr Met Arg Ser Ile Lys Thr Thr Arg
     59              40                  45                  50
     61 gac cga gtg cct aca tat cag tac aac atg aat ttt gaa aag ctg ggc     247
     62 Asp Arg Val Pro Thr Tyr Gln Tyr Asn Met Asn Phe Glu Lys Leu Gly
     63          55                  60                  65
     65 aaa tgc atc ata ata aac aac aag aac ttt gat aaa gtg aca ggt atg     295
```

Missing mandatory <220> feature required with <221>, <222> or <223> features. This error has been indicated in the entire sequence listing. Pleas review and insert <220> where required

RAW SEQUENCE LISTING DATE: 09/22/2000
PATENT APPLICATION: US/09/659,860 TIME: 14:48:08

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

```
66 Lys Cys Ile Ile Ile Asn Asn Lys Asn Phe Asp Lys Val Thr Gly Met
67      70                  75                  80
69 ggc gtt cga aac gga aca gac aaa gat gcc gag gcg ctc ttc aag tgc     343
70 Gly Val Arg Asn Gly Thr Asp Lys Asp Ala Glu Ala Leu Phe Lys Cys
71  85                  90                  95                 100
73 ttc cga agc ctg ggt ttt gac gtg att gtc tat aat gac tgc tct tgt     391
74 Phe Arg Ser Leu Gly Phe Asp Val Ile Val Tyr Asn Asp Cys Ser Cys
75                 105                 110                 115
77 gcc aag atg caa gat ctg ctt aaa aaa gct tct gaa gag gac cat aca     439
78 Ala Lys Met Gln Asp Leu Leu Lys Lys Ala Ser Glu Glu Asp His Thr
79             120                 125                 130
81 aat gcc gcc tgc ttc gcc tgc atc ctc tta agc cat gga gaa gaa aat     487
82 Asn Ala Ala Cys Phe Ala Cys Ile Leu Leu Ser His Gly Glu Glu Asn
83         135                 140                 145
85 gta att tat ggg aaa gat ggt gtc aca cca ata aag gat ttg aca gcc     535
86 Val Ile Tyr Gly Lys Asp Gly Val Thr Pro Ile Lys Asp Leu Thr Ala
87     150                 155                 160
89 cac ttt agg ggg gat aga tgc aaa acc ctt tta gag aaa ccc aaa ctc     583
90 His Phe Arg Gly Asp Arg Cys Lys Thr Leu Leu Glu Lys Pro Lys Leu
91 165                 170                 175                 180
93 ttc ttc att cag gct tgc cga ggg acc gag ctt gat gat ggc atc cag     631
94 Phe Phe Ile Gln Ala Cys Arg Gly Thr Glu Leu Asp Asp Gly Ile Gln
95                 185                 190                 195
97 gcc gac tcg ggg ccc atc aat gac aca gat gct aat cct cga tac aag     679
98 Ala Asp Ser Gly Pro Ile Asn Asp Thr Asp Ala Asn Pro Arg Tyr Lys
99             200                 205                 210
101 atc cca gtg gaa gct gac ttc ctc ttc gcc tat tcc acg gtt cca ggc     727
102 Ile Pro Val Glu Ala Asp Phe Leu Phe Ala Tyr Ser Thr Val Pro Gly
103         215                 220                 225
105 tat tac tcg tgg agg agc cca gga aga ggc tcc tgg ttt gtg caa gcc     775
106 Tyr Tyr Ser Trp Arg Ser Pro Gly Arg Gly Ser Trp Phe Val Gln Ala
107     230                 235                 240
109 ctc tgc tcc atc ctg gag gag cac gga aaa gac ctg gaa atc atg cag     823
110 Leu Cys Ser Ile Leu Glu Glu His Gly Lys Asp Leu Glu Ile Met Gln
111 245                 250                 255                 260
113 atc ctc acc agg gtg aat gac aga gtt gcc agg cac ttt gag tct cag     871
114 Ile Leu Thr Arg Val Asn Asp Arg Val Ala Arg His Phe Glu Ser Gln
115                 265                 270                 275
117 tct gat gac cca cac ttc cat gag aag aag cag atc ccc tgt gtg gtc     919
118 Ser Asp Asp Pro His Phe His Glu Lys Lys Gln Ile Pro Cys Val Val
119             280                 285                 290
121 tcc atg ctc acc aag gaa ctc tac ttc agt caa tag ccatatcagg          965
122 Ser Met Leu Thr Lys Glu Leu Tyr Phe Ser Gln
123         295                 300
125 ggtacattct agctgagaag caatgggtca ctcattaatg aatcacattt ttttatgctc  1025
127 ttgaaatatt cagaaattct ccaggatttt aatttcagga aaatgtattg attcaacagg  1085
129 gaagaaactt tctggtgctg tcttttgttc tctgaatttt cagagacttt tttataatgt  1145
131 tattcatttg gtgactgtgt aactttctct taagattaat tttctctttg tatgtctgtt  1205
133 accttgttaa tagacttaat acatgcaaca gaagtgactt ctggagaaag ctcatggctg  1265
```

RAW SEQUENCE LISTING DATE: 09/22/2000
PATENT APPLICATION: US/09/659,860 TIME: 14:48:08

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

```
     135 tgtccactgc aattggtggt aacagtggta gagtcatgtt tgcacttggc aaaaagaatc  1325
     137 ccaatgtttg acaaaacaca gccaagggga tatttactgc tctttattgc agaatgtggg  1385
     139 tattgagtgt gatttgaatg atttttcatt ggcttagggc agattttcat gcaaaagttc  1445
     141 tcatatgagt tagaggagaa aaagcttaat gattctgata tgtatccatc aggatccagt  1505
     143 ctggaaaaca gaaaccattc taggtgtttc aacagaggga gtttaataca ggaaattgac  1565
     145 ttacatagat gataaaagag aagccaaaca gcaagaagct gttaccacac ccagggctat  1625
     147 gaggataatg ggaagaggtt tggtttcctg tgtccagtag tgggatcatc cagaggagct  1685
     149 ggaaccatgg tgggggctgc ctagtgggag ttaggaccac caatggattg tggaaaatgg  1745
     151 agccatgaca agaacaaagc cactgactga gatggagtga gctgagacag ataagagaat  1805
     153 accttgtctc acctatcctg ccctcacatc ttccaccagc accttactgc ccaggcctat  1865
     155 ctggaagcca cctcaccaag gaccttggaa gagcaaggga cagtgaggca ggagaagaac  1925
     157 aagaaatgga tgtaagcctg gcccataatg tgaacataag taatcactaa tgctcaacaa  1985
     159 tttatccatt caatcattta ttcattgggt tgtcagatag tctatgtatg tgtaaaacaa  2045
     161 tctgttttgg ctttatgtgc aaaatctgtt atagctttaa aatatatctg gaacttttta  2105
     163 gattattcca agccttattt tgagtaaata tttgttactt ttagttctat aagtgaggaa  2165
     165 gagtttatgg caaagatttt tggcactttg ttttcaagat ggtgttatct tttgaattct  2225
     167 tgataaatga ctgtttttt ctgcctaata gtaactggtt aaaaaacaaa tgttcatatt  2285
     169 tattgattaa aaatgtggtt gctt                                       2309
     172 <210> SEQ ID NO: 4
     173 <211> LENGTH: 26
     174 <212> TYPE: DNA
     175 <213> ORGANISM: Artificial Sequence
W--> 177 <220> FEATURE:
     177 <223> OTHER INFORMATION: PCR Primer
     179 <400> SEQUENCE: 4
     180 attggtggta acagtggtag agtcat                                     26
     183 <210> SEQ ID NO: 5
     184 <211> LENGTH: 20
     185 <212> TYPE: DNA
     186 <213> ORGANISM: Artificial Sequence
W--> 188 <220> FEATURE:
     188 <223> OTHER INFORMATION: PCR Primer
     190 <400> SEQUENCE: 5
     191 cccttggctg tgttttgtca                                           20
     194 <210> SEQ ID NO: 6
     195 <211> LENGTH: 27
     196 <212> TYPE: DNA
     197 <213> ORGANISM: Artificial Sequence
W--> 199 <220> FEATURE:
     199 <223> OTHER INFORMATION: PCR Probe
     201 <400> SEQUENCE: 6
     202 ttgcacttgg caaaaagaat cccaatg                                   27
     205 <210> SEQ ID NO: 7
     206 <211> LENGTH: 21
     207 <212> TYPE: DNA
     208 <213> ORGANISM: Artificial Sequence
W--> 210 <220> FEATURE:
     210 <223> OTHER INFORMATION: PCR Primer
     212 <400> SEQUENCE: 7
```

refer to p. 1

RAW SEQUENCE LISTING DATE: 09/22/2000
PATENT APPLICATION: US/09/659,860 TIME: 14:48:08

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

```
213 caacggattt ggtcgtattg g                                              21
216 <210> SEQ ID NO: 8
217 <211> LENGTH: 26
218 <212> TYPE: DNA
219 <213> ORGANISM: Artificial Sequence
W--> 221 <220> FEATURE:
221 <223> OTHER INFORMATION: PCR Primer
223 <400> SEQUENCE: 8
224 ggcaacaata tccactttac cagagt                                          26
227 <210> SEQ ID NO: 9
228 <211> LENGTH: 21
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
W--> 232 <220> FEATURE:
232 <223> OTHER INFORMATION: PCR Probe
234 <400> SEQUENCE: 9
235 cgcctggtca ccagggctgc t                                              21
238 <210> SEQ ID NO: 10
239 <211> LENGTH: 2006
240 <212> TYPE: DNA
241 <213> ORGANISM: Mus musculus
243 <220> FEATURE:
244 <221> NAME/KEY: CDS
245 <222> LOCATION: (474)...(1496)
247 <400> SEQUENCE: 10
248 agctcagtga ggctgatgtg tactgcacat ttaaaaaaaa aatcacagga attttcatac     60
250 aatgaataaa accacaacaa tacatgtaga attggcaggt ggaaaagagc cagcaagggc    120
252 tcaaactaat cactcacttt ccctcttcag catagttcaa ccaacagtag cacactttca    180
254 cctacaaatc ttaaagtagc tccatcaaat ctgcagtttt cacattattg aaaatgtctg    240
256 tcacataggt acaaatttag aatcatcaca ttatattaca tggctattct aggtcatcta    300
258 tagatcagat cttagactac agtgattgaa gttcttcgta cagccatcaa aaagggacac    360
260 atgatcatta cctactgtta gctcacatct aaaggcatga aaaggtttcc ttttttttcaa    420
262 ctgacccaaa cactttaccc caatagtgcc aggttccctc tctgctgctt tga atg       476
263                                                              Met
264                                                               1
266 ttc aca gcc caa gtg ttc tca gag tcc ttt aca aaa act gag ttg ctg      524
267 Phe Thr Ala Gln Val Phe Ser Glu Ser Phe Thr Lys Thr Glu Leu Leu
268               5                  10                  15
270 ccc tcg acc ctt gcg gag gac gga cgc tgc cgt ggg ctc ctg gcc gcc      572
271 Pro Ser Thr Leu Ala Glu Asp Gly Arg Cys Arg Gly Leu Leu Ala Ala
272          20                  25                  30
274 gcc gtg gga acg atg acc gat gat cag gac tgt gct gcg gag ctg gaa      620
275 Ala Val Gly Thr Met Thr Asp Asp Gln Asp Cys Ala Ala Glu Leu Glu
276      35                  40                  45
278 aag gtg gat tct tcc agc gaa gac gga gtt gac gcc aag cca gac cgc      668
279 Lys Val Asp Ser Ser Ser Glu Asp Gly Val Asp Ala Lys Pro Asp Arg
280  50                  55                  60                  65
282 tcc tct atc atc tcc tct att ctc ttg aag aag aag aga aat gcc tct      716
283 Ser Ser Ile Ile Ser Ser Ile Leu Leu Lys Lys Lys Arg Asn Ala Ser
```

Refer to p.1

RAW SEQUENCE LISTING DATE: 09/22/2000
PATENT APPLICATION: US/09/659,860 TIME: 14:48:08

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

```
284                 70                  75                  80
286 gcg ggc ccc gtc agg acc ggc cgg gac cga gtg ccc act tat ctg tac     764
287 Ala Gly Pro Val Arg Thr Gly Arg Asp Arg Val Pro Thr Tyr Leu Tyr
288              85                  90                  95
290 cgc atg gat ttc cag aag atg ggt aaa tgc atc atc ata aac aac aag     812
291 Arg Met Asp Phe Gln Lys Met Gly Lys Cys Ile Ile Ile Asn Asn Lys
292         100                 105                 110
294 aac ttc gac aaa gcg aca ggt atg gac gtc cgg aat ggg acg gac aaa     860
295 Asn Phe Asp Lys Ala Thr Gly Met Asp Val Arg Asn Gly Thr Asp Lys
296     115                 120                 125
298 gat gca ggg gcc ctc ttc aag tgc ttc caa aac ctg ggt ttt gaa gta     908
299 Asp Ala Gly Ala Leu Phe Lys Cys Phe Gln Asn Leu Gly Phe Glu Val
300 130                 135                 140                 145
302 acc gtc cac aat gac tgc tct tgt gca aag atg caa gat ctg ctt aga     956
303 Thr Val His Asn Asp Cys Ser Cys Ala Lys Met Gln Asp Leu Leu Arg
304                 150                 155                 160
306 aaa gcc tct gag gag gac cac agc aac tcg gcc tgc ttc gcc tgc gtc    1004
307 Lys Ala Ser Glu Glu Asp His Ser Asn Ser Ala Cys Phe Ala Cys Val
308             165                 170                 175
310 ctg ctg agc cac ggg gaa gag gac ctg att tac ggg aaa gat ggc gtg    1052
311 Leu Leu Ser His Gly Glu Glu Asp Leu Ile Tyr Gly Lys Asp Gly Val
312         180                 185                 190
314 aca ccc ata aag gat ctg aca gct cat ttt agg gga gac cga tgc aaa    1100
315 Thr Pro Ile Lys Asp Leu Thr Ala His Phe Arg Gly Asp Arg Cys Lys
316     195                 200                 205
318 acc ctg tta gag aaa ccc aaa ctc ttc ttc att cag gca tgc cga ggg    1148
319 Thr Leu Leu Glu Lys Pro Lys Leu Phe Phe Ile Gln Ala Cys Arg Gly
320 210                 215                 220                 225
322 acg gag ctc gac gat gga atc cag gct gac tcg ggg ccc atc aac gac    1196
323 Thr Glu Leu Asp Asp Gly Ile Gln Ala Asp Ser Gly Pro Ile Asn Asp
324                 230                 235                 240
326 att gac gct aat ccc cgc aac aag atc ccg gtg gaa gcc gac ttc ctc    1244
327 Ile Asp Ala Asn Pro Arg Asn Lys Ile Pro Val Glu Ala Asp Phe Leu
328             245                 250                 255
330 ttt gct tac tcc acg gtt cca ggt tat tac tca tgg agg aac cca ggg    1292
331 Phe Ala Tyr Ser Thr Val Pro Gly Tyr Tyr Ser Trp Arg Asn Pro Gly
332         260                 265                 270
334 aaa ggc tcc tgg ttt gtg cag gcc ctc tgc tcc atc ctg aat gag cat    1340
335 Lys Gly Ser Trp Phe Val Gln Ala Leu Cys Ser Ile Leu Asn Glu His
336     275                 280                 285
338 ggc aag gac ctc gag atc atg cag atc ctg acc agg gtg aac gac agg    1388
339 Gly Lys Asp Leu Glu Ile Met Gln Ile Leu Thr Arg Val Asn Asp Arg
340 290                 295                 300                 305
342 gtg gcc agg cac ttc gag tcc cag tct gat gat cca cgc ttc aac gag    1436
343 Val Ala Arg His Phe Glu Ser Gln Ser Asp Asp Pro Arg Phe Asn Glu
344                 310                 315                 320
346 aag aag cag atc ccg tgt atg gtg tcc atg ctc acc aaa gag ctg tac    1484
347 Lys Lys Gln Ile Pro Cys Met Val Ser Met Leu Thr Lys Glu Leu Tyr
348             325                 330                 335
```

VERIFICATION SUMMARY DATE: 09/22/2000
PATENT APPLICATION: US/09/659,860 TIME: 14:48:09

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:18 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:29 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:177 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:188 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:199 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:210 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:221 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:232 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:376 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:387 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:398 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:409 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:420 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:431 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:873 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:884 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:895 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:906 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:917 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:928 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:939 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:950 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:961 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:972 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:983 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:994 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1005 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1016 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1027 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1038 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1049 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1060 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1071 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1082 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1093 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1104 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1115 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1126 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1137 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1148 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1159 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1170 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1181 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1192 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1203 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1214 M:258 W: Mandatory Feature missing, <220> FEATURE:

VERIFICATION SUMMARY DATE: 09/22/2000
PATENT APPLICATION: US/09/659,860 TIME: 14:48:09

Input Set : A:\RTS-0201_Seq_ASCII.txt
Output Set: N:\CRF3\09222000\I659860.raw

L:1225 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1236 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1247 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:1258 M:258 W: Mandatory Feature missing, <220> FEATURE: